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Pro Ser Asp Val Gln Pro Pro Gly Asp Pro Phe Gln Val Asn Phe Glu
            260             265             270

gaa caa aac aat cct caa ata ctc caa aac tca gtt gtt ttt gga aca        864
Glu Gln Asn Asn Pro Gln Ile Leu Gln Asn Ser Val Val Phe Gly Thr
    275             280             285
```

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```
tca gcc cag gaa gtg gta aag gaa att cgt ttc aga att gag cag aaa    912
Ser Ala Gln Glu Val Val Lys Glu Ile Arg Phe Arg Ile Glu Gln Lys
    290                 295                 300

aca aca ctg aca gcc agt gca ggt gtt cgg ata tct agt ttt ccc aat    960
Thr Thr Leu Thr Ala Ser Ala Gly Val Arg Ile Ser Ser Phe Pro Asn
305                 310                 315                 320

gaa gag gac agg aaa cac caa caa agg agc att att ggc ttt tta cag   1008
Glu Glu Asp Arg Lys His Gln Gln Arg Ser Ile Ile Gly Phe Leu Gln
                325                 330                 335

gct gga aac caa gcc ctg tca gcc act gag tgt aca tta gag aaa act   1056
Ala Gly Asn Gln Ala Leu Ser Ala Thr Glu Cys Thr Leu Glu Lys Thr
                340                 345                 350

gac aaa gat aag ttt gta aaa cct cta gaa atg tct cat aag aag agt   1104
Asp Lys Asp Lys Phe Val Lys Pro Leu Glu Met Ser His Lys Lys Ser
                355                 360                 365

ttc ttt gat aaa aaa cga tca gaa agg aaa tgg agt cac caa gat aca   1152
Phe Phe Asp Lys Lys Arg Ser Glu Arg Lys Trp Ser His Gln Asp Thr
    370                 375                 380

ttt aaa tgt gaa gcc gtg aat aaa caa agt ttc cag aca tca caa cca   1200
Phe Lys Cys Glu Ala Val Asn Lys Gln Ser Phe Gln Thr Ser Gln Pro
385                 390                 395                 400

ttc caa gtt tta aag aag aag atg aat gag aat ttg gaa ata tca gag   1248
Phe Gln Val Leu Lys Lys Lys Met Asn Glu Asn Leu Glu Ile Ser Glu
                405                 410                 415

aat tca gat gac tgt cag ata ctt acc tgt cct gtt tgc ttt agg gct   1296
Asn Ser Asp Asp Cys Gln Ile Leu Thr Cys Pro Val Cys Phe Arg Ala
                420                 425                 430

caa ggg tgc atc agt ctg gaa gcc ttg aat aaa cat gta gat gaa tgt   1344
Gln Gly Cys Ile Ser Leu Glu Ala Leu Asn Lys His Val Asp Glu Cys
                435                 440                 445

ctt gat gga cct tca atc agt gaa aac ttt aaa atg ttc tcg tgt tca   1392
Leu Asp Gly Pro Ser Ile Ser Glu Asn Phe Lys Met Phe Ser Cys Ser
    450                 455                 460

cat gtt tct gct acc aaa gtt aac aag aaa gaa aat gtt cct gct tct   1440
His Val Ser Ala Thr Lys Val Asn Lys Lys Glu Asn Val Pro Ala Ser
465                 470                 475                 480

tca ctt tgt gag aag caa gat tat gaa gcc cat cca aaa att aaa gaa   1488
Ser Leu Cys Glu Lys Gln Asp Tyr Glu Ala His Pro Lys Ile Lys Glu
                485                 490                 495

ata tct tca gta gat tgt ata gct tta gta gat act ata gat aac tca   1536
Ile Ser Ser Val Asp Cys Ile Ala Leu Val Asp Thr Ile Asp Asn Ser
                500                 505                 510

tct aaa gca gaa agc ata gat gct tta agt aat aag cat agc aag gaa   1584
```

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```
Ser Lys Ala Glu Ser Ile Asp Ala Leu Ser Asn Lys His Ser Lys Glu
        515                 520                 525

gaa tgt tct agt ctc cca agc aag tct ttt aat att gaa cac tgt cat      1632
Glu Cys Ser Ser Leu Pro Ser Lys Ser Phe Asn Ile Glu His Cys His
        530                 535                 540

cag aat tct tct tct act gtt tca ttg gaa aac gaa gat gtt gga tca      1680
Gln Asn Ser Ser Ser Thr Val Ser Leu Glu Asn Glu Asp Val Gly Ser
545                 550                 555                 560

ttt aga caa gaa tac cgc cag cct tac tta tgt gaa gtg aaa aca ggc      1728
Phe Arg Gln Glu Tyr Arg Gln Pro Tyr Leu Cys Glu Val Lys Thr Gly
                565                 570                 575

caa gct cta gtt tgt cct gtt tgt aac gta gaa caa aag act tca gat      1776
Gln Ala Leu Val Cys Pro Val Cys Asn Val Glu Gln Lys Thr Ser Asp
                580                 585                 590

cta acc ctg ttc aat gtg cat gtg gat gtt tgc tta aat aaa agt ttt      1824
Leu Thr Leu Phe Asn Val His Val Asp Val Cys Leu Asn Lys Ser Phe
                595                 600                 605

atc caa gaa tta aga aag gat aaa ttt aac cca gtt aat caa ccc aaa      1872
Ile Gln Glu Leu Arg Lys Asp Lys Phe Asn Pro Val Asn Gln Pro Lys
        610                 615                 620

gaa agc tcc aga agt act ggt agc tca agt gga gta cag aag gct gta      1920
Glu Ser Ser Arg Ser Thr Gly Ser Ser Ser Gly Val Gln Lys Ala Val
625                 630                 635                 640

aca aga aca aaa agg cca gga ttg atg aca aag tac tca aca tca aag      1968
Thr Arg Thr Lys Arg Pro Gly Leu Met Thr Lys Tyr Ser Thr Ser Lys
                645                 650                 655

aaa ata aaa cca aac aat ccc aaa cat acc ctt gat ata ttt ttt aag      2016
Lys Ile Lys Pro Asn Asn Pro Lys His Thr Leu Asp Ile Phe Phe Lys
                660                 665                 670

taagtcgacc                                                           2026


<210> 31
<211> 672
<212> PRT
<213> Homo sapiens

<400> 31
Met Asp Ser Thr Lys Glu Lys Cys Asp Ser Tyr Lys Asp Asp Leu Leu
  1                 5                 10                  15

Leu Arg Met Gly Leu Asn Asp Asn Lys Ala Gly Met Glu Gly Leu Asp
                20                  25                  30

Lys Glu Lys Ile Asn Lys Ile Ile Met Glu Ala Thr Lys Gly Ser Arg
        35                  40                  45
```

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```
Phe Tyr Gly Asn Glu Leu Lys Lys Glu Lys Gln Val Asn Gln Arg Ile
    50                  55                  60

Glu Asn Met Met Gln Gln Lys Ala Gln Ile Thr Ser Gln Gln Leu Arg
 65              70                  75                      80

Lys Ala Gln Leu Gln Val Asp Arg Phe Ala Met Glu Leu Glu Gln Ser
                85                  90                  95

Arg Asn Leu Ser Asn Thr Ile Val His Ile Asp Met Asp Ala Phe Tyr
            100                 105             110

Ala Ala Val Glu Met Arg Asp Asn Pro Glu Leu Lys Asp Lys Pro Ile
        115                 120             125

Ala Val Gly Ser Met Ser Met Leu Ser Thr Ser Asn Tyr His Ala Arg
    130                 135                 140

Arg Phe Gly Val Arg Ala Ala Met Pro Gly Phe Ile Ala Lys Arg Leu
145             150                 155                 160

Cys Pro Gln Leu Ile Ile Val Pro Pro Asn Phe Asp Lys Tyr Arg Ala
                165                 170                 175

Val Ser Lys Glu Val Lys Glu Ile Leu Ala Asp Tyr Asp Pro Asn Phe
            180                 185             190

Met Ala Met Ser Leu Asp Glu Ala Tyr Leu Asn Ile Thr Lys His Leu
        195                 200                 205

Glu Glu Arg Gln Asn Trp Pro Glu Asp Lys Arg Arg Tyr Phe Ile Lys
    210             215                 220

Met Gly Ser Ser Val Glu Asn Asp Asn Pro Gly Lys Glu Val Asn Lys
225             230                 235                     240

Leu Ser Glu His Glu Arg Ser Ile Ser Pro Leu Leu Phe Glu Glu Ser
            245                 250                 255

Pro Ser Asp Val Gln Pro Pro Gly Asp Pro Phe Gln Val Asn Phe Glu
            260                 265                 270

Glu Gln Asn Asn Pro Gln Ile Leu Gln Asn Ser Val Val Phe Gly Thr
    275                 280                 285

Ser Ala Gln Glu Val Val Lys Glu Ile Arg Phe Arg Ile Glu Gln Lys
    290                 295                 300

Thr Thr Leu Thr Ala Ser Ala Gly Val Arg Ile Ser Ser Phe Pro Asn
305             310                 315                     320

Glu Glu Asp Arg Lys His Gln Gln Arg Ser Ile Ile Gly Phe Leu Gln
            325                 330                 335

Ala Gly Asn Gln Ala Leu Ser Ala Thr Glu Cys Thr Leu Glu Lys Thr
            340                 345                 350
```

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Asp Lys Asp Lys Phe Val Lys Pro Leu Glu Met Ser His Lys Lys Ser
   355   360   365

Phe Phe Asp Lys Lys Arg Ser Glu Arg Lys Trp Ser His Gln Asp Thr
   370   375   380

Phe Lys Cys Glu Ala Val Asn Lys Gln Ser Phe Gln Thr Ser Gln Pro
385   390   395   400

Phe Gln Val Leu Lys Lys Lys Met Asn Glu Asn Leu Glu Ile Ser Glu
               405   410   415

Asn Ser Asp Asp Cys Gln Ile Leu Thr Cys Pro Val Cys Phe Arg Ala
               420   425   430

Gln Gly Cys Ile Ser Leu Glu Ala Leu Asn Lys His Val Asp Glu Cys
           435   440   445

Leu Asp Gly Pro Ser Ile Ser Glu Asn Phe Lys Met Phe Ser Cys Ser
   450   455   460

His Val Ser Ala Thr Lys Val Asn Lys Lys Glu Asn Val Pro Ala Ser
465   470   475   480

Ser Leu Cys Glu Lys Gln Asp Tyr Glu Ala His Pro Lys Ile Lys Glu
               485   490   495

Ile Ser Ser Val Asp Cys Ile Ala Leu Val Asp Thr Ile Asp Asn Ser
               500   505   510

Ser Lys Ala Glu Ser Ile Asp Ala Leu Ser Asn Lys His Ser Lys Glu
           515   520   525

Glu Cys Ser Ser Leu Pro Ser Lys Ser Phe Asn Ile Glu His Cys His
   530   535   540

Gln Asn Ser Ser Ser Thr Val Ser Leu Glu Asn Glu Asp Val Gly Ser
545   550   555   560

Phe Arg Gln Glu Tyr Arg Gln Pro Tyr Leu Cys Glu Val Lys Thr Gly
               565   570   575

Gln Ala Leu Val Cys Pro Val Cys Asn Val Glu Gln Lys Thr Ser Asp
           580   585   590

Leu Thr Leu Phe Asn Val His Val Asp Val Cys Leu Asn Lys Ser Phe
           595   600   605

Ile Gln Glu Leu Arg Lys Asp Lys Phe Asn Pro Val Asn Gln Pro Lys
   610   615   620

Glu Ser Ser Arg Ser Thr Gly Ser Ser Ser Gly Val Gln Lys Ala Val
625   630   635   640

Thr Arg Thr Lys Arg Pro Gly Leu Met Thr Lys Tyr Ser Thr Ser Lys
               645   650   655

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Lys Ile Lys Pro Asn Asn Pro Lys His Thr Leu Asp Ile Phe Phe Lys
            660   665   670

<210> 32
<211> 1335
<212> DNA
<213> Homo sapiens

<400> 32
tgagagagct ttccgctgaa gatgacgggc ctgctttcca gggcggcttg tcgaaagccc 60

gggagcatct ggccgcttcc gcctcaacca tgggctgggg ttttgtgagc tactagtgcc 120

aagggttttc tttccaccag accaccgctg taaatctcga gggtcttact cattagaagt 180

tagaattcac atttgacgtt taaaggaaga atttccttag taccttctca caagcacgca 240

cttcgcattt ttagatttct agagtttgct ttgtagaaag taattttgag gttgtcagag 300

aataaatgac gttagaaagg tttttaaagt aaaacaagaa tgtgagatga tagcctggga 360

ttttctcttg gttgtaaatg aatatcttac tgagaaccac gttaaccatg cctgcccctc 420

aaagatagga aaggttggat atatagaaac tttctcgtat tagaaatacc gaagtgcagt 480

ggttttgtgt gtacaaggga ttaggcaata ggaggctatt tttgttttaa gactagggtt 540

gaattagcag aaagaccaat agaagatcta caactcttg tcagttgtca aggataactt 600

tgattatgag actttgactt tgtagcttca gtaatttcct ctcgttagct attttaatat 660

agtcgatttc cttgtaattg ccaagagtaa aatttgttat taaaccttag aaagagtact 720

ttcttactac aaggatggga cgataggagc gaaatttcga gtctaaggga aaacgctggc 780

cgagtgtggt ggctcacgcc tgtaatccca gcacttcggg aggccgaggt gggtggatca 840

cctgaggccg ggagtttgag accagcctgg caacaagat ttttcttcat ccctttactt 900

tgagtctgtg gatgtcattg catgtgatat gggtctcctg aagacagcat accattggat 960

tttgcttctt tatccaagtt atcattctgt cttttaattg gggtgtgcat tcaagataag 1020

tttataccat ggatagcaca aaggagaagt gtgacagtta caaagatgat cttctgctta 1080

ggatgggact taatgataat aaagcaggaa tggaaggatt agataaagag aaaattaaca 1140

aaattataat ggaagccacg aaggggtcca gattttatgg aaatgagctc aagaaagaaa 1200

agcaagtcaa ccaacgaatt gaaaatatga tgcaacaaaa agctcaaatc accagccaac 1260

agctaagaaa agcacaatta caggttgaca gatttgcaat ggaattagaa caaagccgaa 1320

atttgagcaa tacca   1335

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<210> 33
<211> 105
<212> DNA
<213> Mus musculus

<400> 33
gggagcgtcg cgagccgccg ggaggggccc ggggcggggt ggagggagga tgggaggacg 60

gaggggaggg agctgagaga ggagggaggg taaatagtgg acccg 105

<210> 34
<211> 105
<212> DNA
<213> Mus musculus

<400> 34
cgggtccact atttaccctc cctcctctct cagctccctc ccctccgtcc tcccatcctc 60

cctccacccc gccccgggcc cctcccggcg gctcgcgacg ctccc 105

<210> 35
<211> 140
<212> DNA
<213> Homo sapiens

<400> 35
ccctgcttat atagatgacc ccctccccga gactctgaca gacccaggtc acaggcagtc 60

ctcacctgct cctgacaccc ccggcccctc agtgctgctc tctctagcca ccgagctgaa 120

gtactgagga gcccctacct 140

<210> 36
<211> 140
<212> DNA
<213> Homo sapiens

<400> 36
aggtaggggc tcctcagtac ttcagctcgg tggctagaga gagcagcact gaggggccgg 60

gggtgtcagg agcaggtgag gactgcctgt gacctgggtc tgtcagagtc tcggggaggg 120

ggtcatctat ataagcaggg 140

<210> 37
<211> 37
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-GPBP-6c

```
<400> 37
ctcgctcgcc cagggaagga aaagggaaaa gaaggga                              37


<210> 38
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-GPBP-14c

<400> 38
ctgcctggcc cactatttac c                                              21


<210> 39
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-GPBP-18m

<400> 39
ggcatggtta acgtggttct c                                              21


<210> 40
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-XbaG/Bpro1m

<400> 40
gactctagag ggttcgggag gaggatcccg                                     30


<210> 41
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-XbaG/Bpro1c

<400> 41
gactctagac tggcccacta tttaccctcc                                     30


<210> 42
<211> 30
<212> DNA
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Primer ON-SP1Del

<400> 42
cgccgggagg gggacgtagt gggggagaat   30

<210> 43
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-TATADel

<400> 43
caggggaggg gaggggtggg ccagtctaga   30

<210> 44
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-DIN2c

<400> 44
ggattattgc acttgccttc ac   22

<210> 45
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-DIN5'm

<400> 45
aaaggatcca tggatagcac aaaggag   27

<210> 46
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-DIN-THc

<400> 46
aaaaaagtcg acttacttaa aaaatatatc aagggt   36

<210> 47
<211> 21

```
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-DINB1-R2

<400> 47
tggtattgct caaatttcgg c                                          21


<210> 48
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-GPBP-39c

<400> 48
tgagagagct ttccgctg                                              18


<210> 49
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-LMPTAP1m

<400> 49
atgtctagat gtgtagggca gatctgccc                                  29


<210> 50
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-LMPTAP1c

<400> 50
atgtctagac tggtgcccaa ttttctcca                                  29


<210> 51
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-HSP1m

<400> 51
atgtctagat aagccggccg gagagggct                                  29
```

<210> 52
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-HSP1c

<400> 52
atgtctagac gcggcaccgc gtgtgcagg   29

<210> 53
<211> 31
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-SA3A4m

<400> 53
gactctagag ggttaaggag gtgatgctcc c   31

<210> 54
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-SA3A4c

<400> 54
gactctagat ggccactccc tccaccctgc gc   32

<210> 55
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-INGA3A4m

<400> 55
gactctagac acccaggctt tttggttgtg gc   32

<210> 56
<211> 31
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-INGA3A4c

```
<400> 56
gactctagaa agcggggcct cccgcagacg c                                    31


<210> 57
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-S2A3A4m

<400> 57
atgtctagat aggcactgga caagccccc                                      29


<210> 58
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-S2A3A4c

<400> 58
atgtctagag ggctagtggc gaggctgag                                      29


<210> 59
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-IDH-F1

<400> 59
cacagagggc gagtacagca                                                20


<210> 60
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-IDH-R1

<400> 60
tgatcttcag gctctccacc a                                              21


<210> 61
<211> 20
<212> DNA
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Primer ON-TRAPD-F1

<400> 61
gggtccagaa catggctctc   20

<210> 62
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-TRAPD-R1

<400> 62
acatcctggc ctcgagtgac   20

<210> 63
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-LMP2-F2

<400> 63
gcagcatata agccaggcat g   21

<210> 64
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-LMP2-R2

<400> 64
tggccagagc aatagcgtct   20

<210> 65
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-TAP1-F2

<400> 65
gccgcctcac tgactggat   19

<210> 66
<211> 21

24

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-TAP1-R2

<400> 66
tcgagtgaag gtatcggctg a   21

<210> 67
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-DHFR-F1

<400> 67
cctgtggagg aggaggtgg   19

<210> 68
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-DHFR-R1

<400> 68
ccgattcttc cagtctacgg g   21

<210> 69
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-MSH3-F1

<400> 69
tgggtaaagg ttggaagcac a   21

<210> 70
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-MSH3-R1

<400> 70
aaaaggagag tgaaagcggc t   21

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<210> 71
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-HO3-F2

<400> 71
gagctgttgt ccctccgct                                          19


<210> 72
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-HO3-R2

<400> 72
ggccagataa cgagcaaagg                                         20


<210> 73
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-HARS-F2

<400> 73
aggtggcgaa actcctgaaa c                                       21


<210> 74
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-HARS-R2

<400> 74
tgctttcatc aggacccagc                                         20


<210> 75
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-Hsp10-F1
```

```
<400> 75
ggagggagta atggcaggac a                                              21


<210> 76
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-Hsp10-R1

<400> 76
agcagcactc ctttcaacca a                                              21


<210> 77
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-Hsp60-F1

<400> 77
gcctttggtc ataatcgctg a                                              21


<210> 78
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-Hsp60-R1

<400> 78
tgccacaacc tgaagaccaa c                                              21


<210> 79
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A1-F1

<400> 79
gctctacgtg caaggcaatg a                                              21


<210> 80
<211> 20
<212> DNA
<213> Artificial Sequence
```

27

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A1-R1

<400> 80
attgtgctga acttgcgcag   20

<210> 81
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A2-F1

<400> 81
gaaaagggtg acgtagggca   20

<210> 82
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A2-R1

<400> 82
ggtgtctgat ggaatcccgt t   21

<210> 83
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-GP-F1

<400> 83
ggagacagtg gatcacctgc a   21

<210> 84
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-GP-R1

<400> 84
tgctgtggtt tgactgtgtc g   21

<210> 85
<211> 21

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A4-F1

<400> 85
cttgccttcc cgtatttagc a   21

<210> 86
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A4-R1

<400> 86
ggatctgtcg tttctctggg c   21

<210> 87
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A5-F1

<400> 87
catcgaatgt catgggaggg   20

<210> 88
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A5-R1

<400> 88
agttgccagc caaaagctgt a   21

<210> 89
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A6-F1

<400> 89
tttgggctag actaccggac a   21

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<210> 90
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-COL4A6-R1

<400> 90
tctctatgga cccgagggct                                              20


<210> 91
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-GPBP-F1

<400> 91
ctgaatccag cttgcgtcg                                               19


<210> 92
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-GPBP-R1

<400> 92
gcagagtagc cacttgctcc                                              20


<210> 93
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-DinB1-F3

<400> 93
gccccccaac tttgacaaat                                              20


<210> 94
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-DinB1-R3
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<400> 94
gcttcatcaa gactcatggc c   21

<210> 95
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-hGAPDH-F1

<400> 95
gaaggtgaag gtcggagtc   19

<210> 96
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-hGAPDH-R1

<400> 96
gaagatggtg atgggatttc   20

<210> 97
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-GPBP-26-1F

<400> 97
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<210> 98
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-mGPBP-26-1R

<400> 98
ccatttcttc aaccttttgt acaa   24

<210> 99
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
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<400> 99
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<210> 100
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-huDINB-76-F1

<400> 100
ccagtgcagg tgttcggata   20

<210> 101
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-huDINB-76-R1

<400> 101
tttccagcct gtaaaaagcc a   21

<210> 102
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer ON-hGPBP-26-1R

<400> 102
ccatctcttc aacctttgg aca   23